

Figure 1A

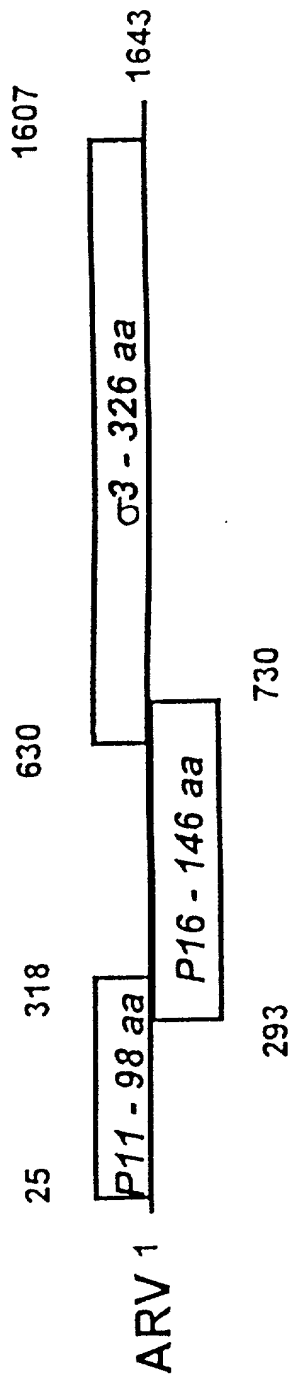


Figure 1B

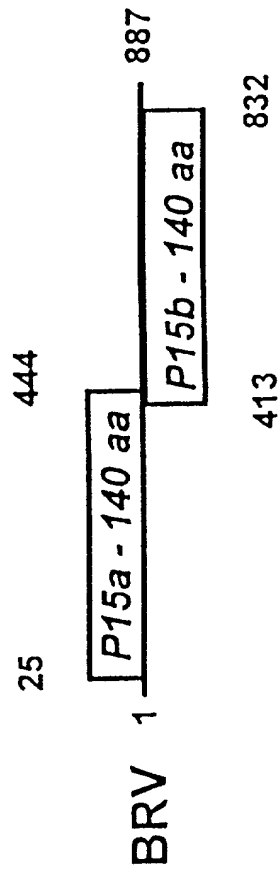


Figure 1C

	*	* *	transmembrane
ARV-138	1	MLRMPPGSCN GATAIFGNVH COAQNTAGG DLOATSSIIA YWPYLAAGGG FLLIIIFI AIF	61
ARV-176	1	MLRMPPGSCN GATAVFGNVH COAQNTAGG DLOATSSIIA YWPYLAAGGG FLLIIVII FALL	61
NBV	1	. . . M SSDCA KIVSVFGSVH CQSSKNSAGG DLOATSVFTT WPHFAIGGG IIVVILLGLF	57
Consensus		- - - - - C - - - - - FG-VH CQ - - - N-AGG DLOATS - - - - - YWP - - A-GGG - - - - -	
		** + + + +	
ARV-138	62	YCCKAKVKAD AARSVFHREL VALSSGKHNA MAPPYDV.	98
ARV-176	62	YCCKAKVKAD AARSVFHREL VALSSGKHNA MAPPYDV.	98
NBV	58	YCCYLKWKTS QVKHTYRREL IALTRSHVHS TPSGISYV	95
consensus		YCC--K-K-- -- -- -- -- REL -AL- -- -- -- --	

Figure 2

transmembrane

+ ++ + ++++

1 MGQRHSIVQP PAPPNAFVE IVSSSTGIII AVGIFAFIFS FLYKLLQWYN RSKNKKRKE 60

61 QIREQIELGL LSYGAGVASL PLLNVIAHNP GSVISATPIY KGPCTGVPNS RLLQITSGTA 120

121 EENTRILNHD GRNPDGSINV

Figure 3